

# Resource Summary Report

Generated by [SPARC SAWG](#) on Sep 17, 2026

## NuChart

RRID:SCR\_004703

Type: Tool

### Proper Citation

NuChart (RRID:SCR\_004703)

### Resource Information

**URL:** <http://www.plosone.org/article/info%3Adoi%2F10.1371%2Fjournal.pone.0075146>

**Proper Citation:** NuChart (RRID:SCR\_004703)

**Description:** An R Package to Study Gene Spatial Neighbourhoods with Multi-Omics Annotations.

**Abbreviations:** NuChart

**Resource Type:** software resource

**Resource Name:** NuChart

**Resource ID:** SCR\_004703

**Alternate IDs:** OMICS\_00525

**Record Creation Time:** 20220129T080226+0000

**Record Last Update:** 20260912T195615+0000

### Ratings and Alerts

No rating or validation information has been found for NuChart.

No alerts have been found for NuChart.

### Data and Source Information

**Source:** [SciCrunch Registry](#)

### Usage and Citation Metrics

We found 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [SPARC SAWG](#) .

D'Agostino D, et al. (2021) Advantages of using graph databases to explore chromatin conformation capture experiments. BMC bioinformatics, 22(Suppl 2), 43.

Tordini F, et al. (2016) The Genome Conformation As an Integrator of Multi-Omic Data: The Example of Damage Spreading in Cancer. Frontiers in genetics, 7, 194.